

6555344

FIG.1

SREB 1 MANASEPGGSGGGEAAALG---DKLATLSLLLCVSLAGN 36
SREB 2 MANYSHAADNILONLSP--LTAFKLTSLGFIIGVSVVGN 38
SREB 3 MANTTGEPEEVSGALSEP SASAYVKEVLGLIMCVSLAGN 40

SREB 1 VLFALLIVRERSLHRAPYYLLDLCLADGLRALACLPAVM 76
SREB 2 LLISILLVKDKTLHRAPYYFLDLCCSDILRSAICFPFVF 78
SREB 3 AILSELLVLKERALHKAPYYFLDLCLADGIRSAVCFPFVL 80

SREB 1 LAARRAAAAAGAPP GALGCKLLAFLAALFCFHAAFLLLGV 116
SREB 2 NSVKNGSTWTY---GTLTCKVIAFLGVLSCFHTAFMLFCI 115
SREB 3 ASVRHGSSTWF---SALSCKIVAFMAVLFCEHAAFMFLFCI 117

SREB 1 GVTRYLAIAHHRFYAERLAGWPCAAMLVCAAWALALAAAF 156
SREB 2 SVTRYLAIAHHRFYTKRLTFWTCLAV-ICMVWTL SVAMAF 154
SREB 3 SVTRYMAIAHHRFYAKRMTLWTCAAV-ICMAWTL SVAMAF 156

SREB 1 PPVLDGGG---DDEDA PCAL EQRPDGAPGALGFLLLLAVV 193
SREB 2 PPVLDVGTYSFIREEDQCTFOHRSFRANDSLGFMLLLALI 194
SREB 3 PPVFDVGTYKFIREEDQCI FEHRYFKANDTLGFMLMLAVL 196

SREB 1 VGATHLVYLRLLFFIHDRRKMRPARLVPAVSHDWTFHGPG 233
SREB 2 LLATQLVYLKLIFFVHDRRKMKPVQFVAAVSONWTFHGPG 234
SREB 3 MAATHAVYGKLLLFEYRHRKMKPVQMVPALSONWTFHGPG 236

SREB 1 ATGOAAANWTAGFGRGPTPPALVGIRPAGPGRGARLLVL 273
SREB 2 ASGOAAANWL AGFGRGPTPTLLGIRQNA NT TGRRLLVL 274
SREB 3 ATGOAAANWIAGFGRGPMPTLLGIRONGHAA SRR-LLGM 275

SREB 1 EEFKTEKRLCKMFYAVTLLFLLWGPYVVASYLRLVLRPG 313
SREB 2 DEFKMEKRISRMFYIMTELFLLWGPYLVACYWRVFARGP 314
SREB 3 DEVKGEKQLGRMFYAITLLFLLWSPYIVACYWRVFVKAC 315

SREB 1 AVPQA YLTASVWLTFAQAGINP VVCFLFNREL RD CFRAQF 353
SREB 2 VVPGGFLTA AVWMSFAQAGINP FVCIFSNRELRCFSTTL 354
SREB 3 AVPHRYLAT AVWMSFAQA AVNP IVCFLLNKDLKKCLRT HA 355

SREB 1 EC CQSPRTTQATHP--CDLKGIGL 376
SREB 2 LYCRKS---RLPREPYC-----VI 371
SREB 3 E- CWGTGGAPAPREPYC-----VM 374

FIG.2

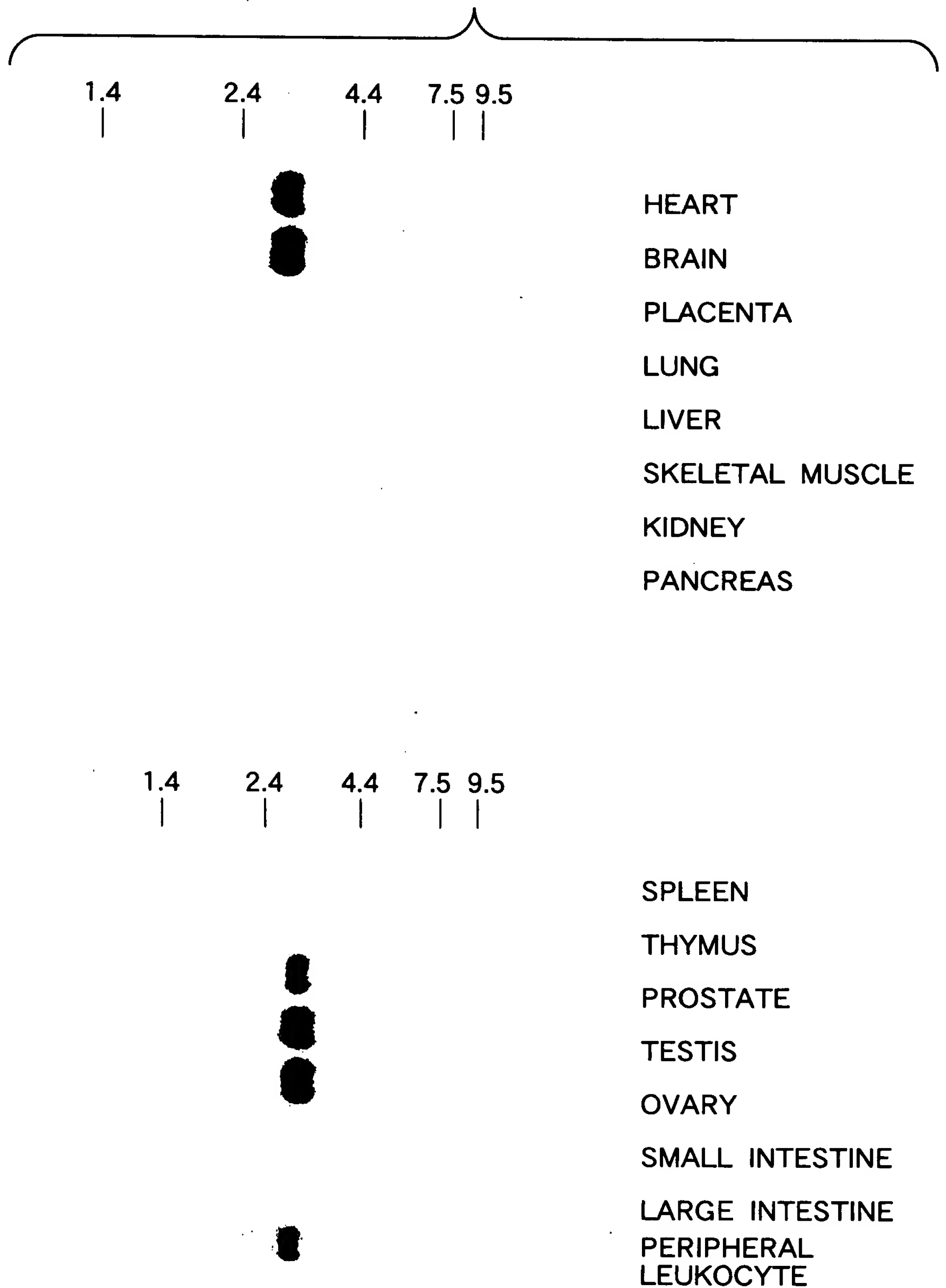


FIG.3

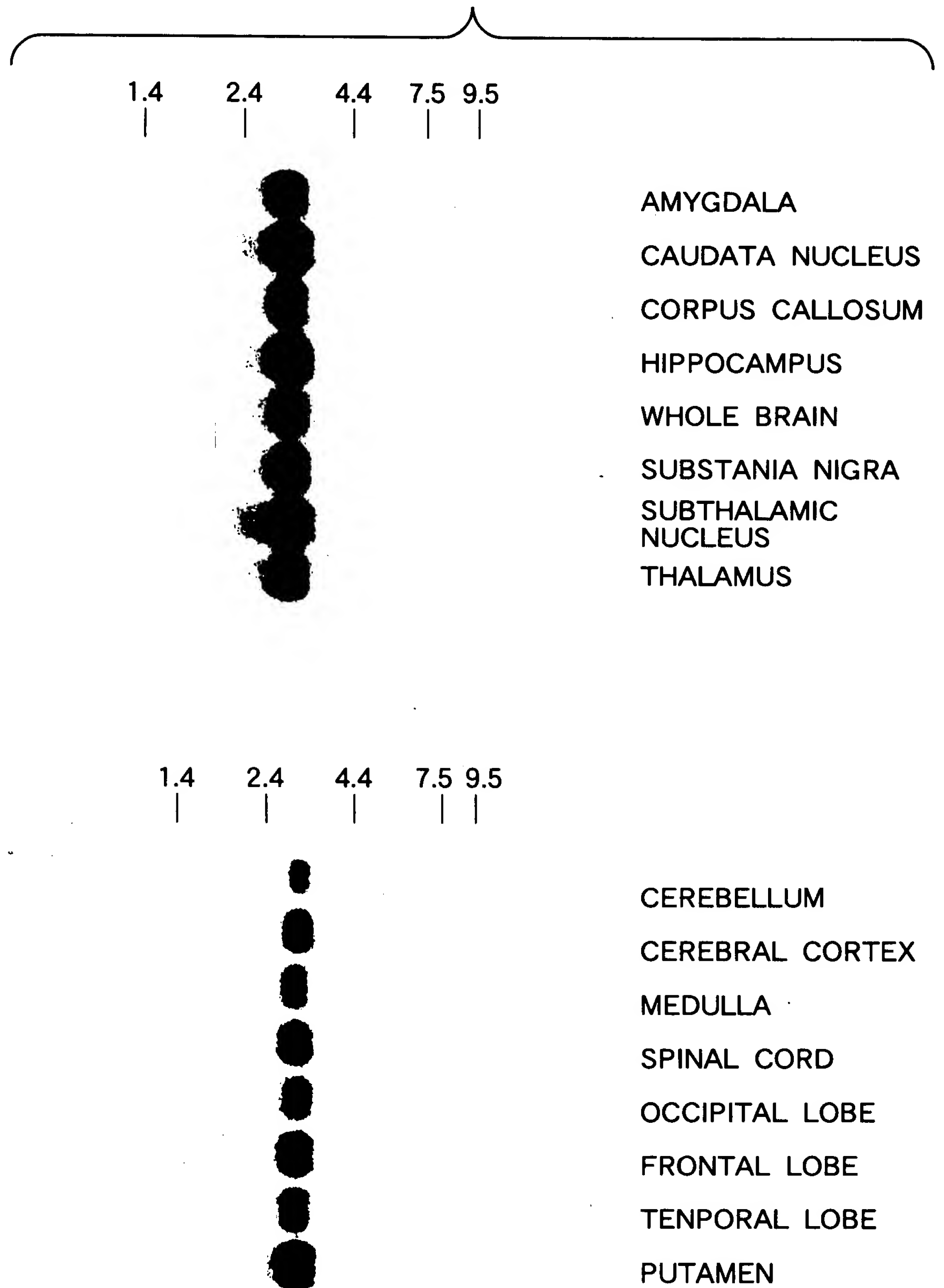


FIG.4

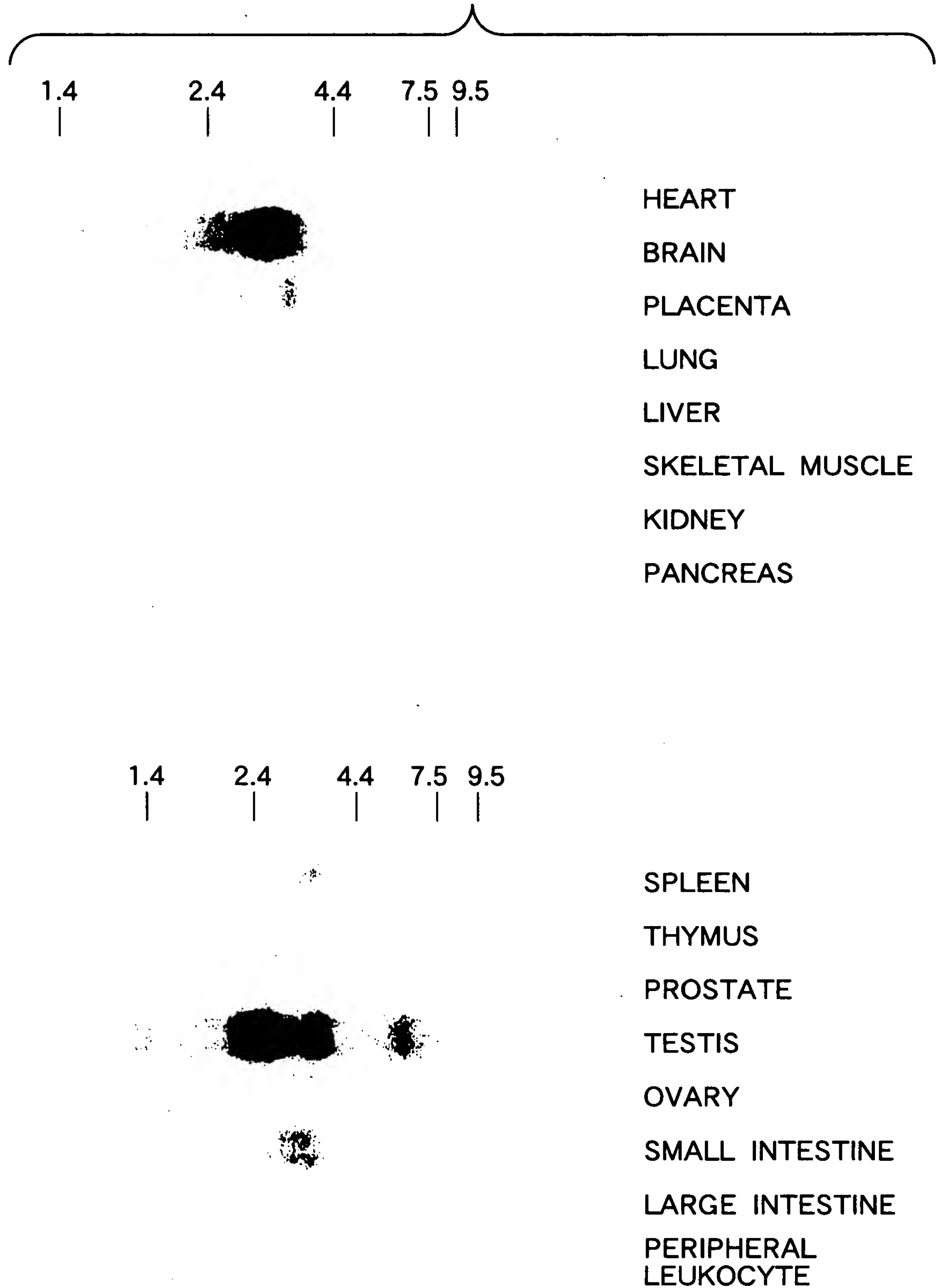


FIG.5

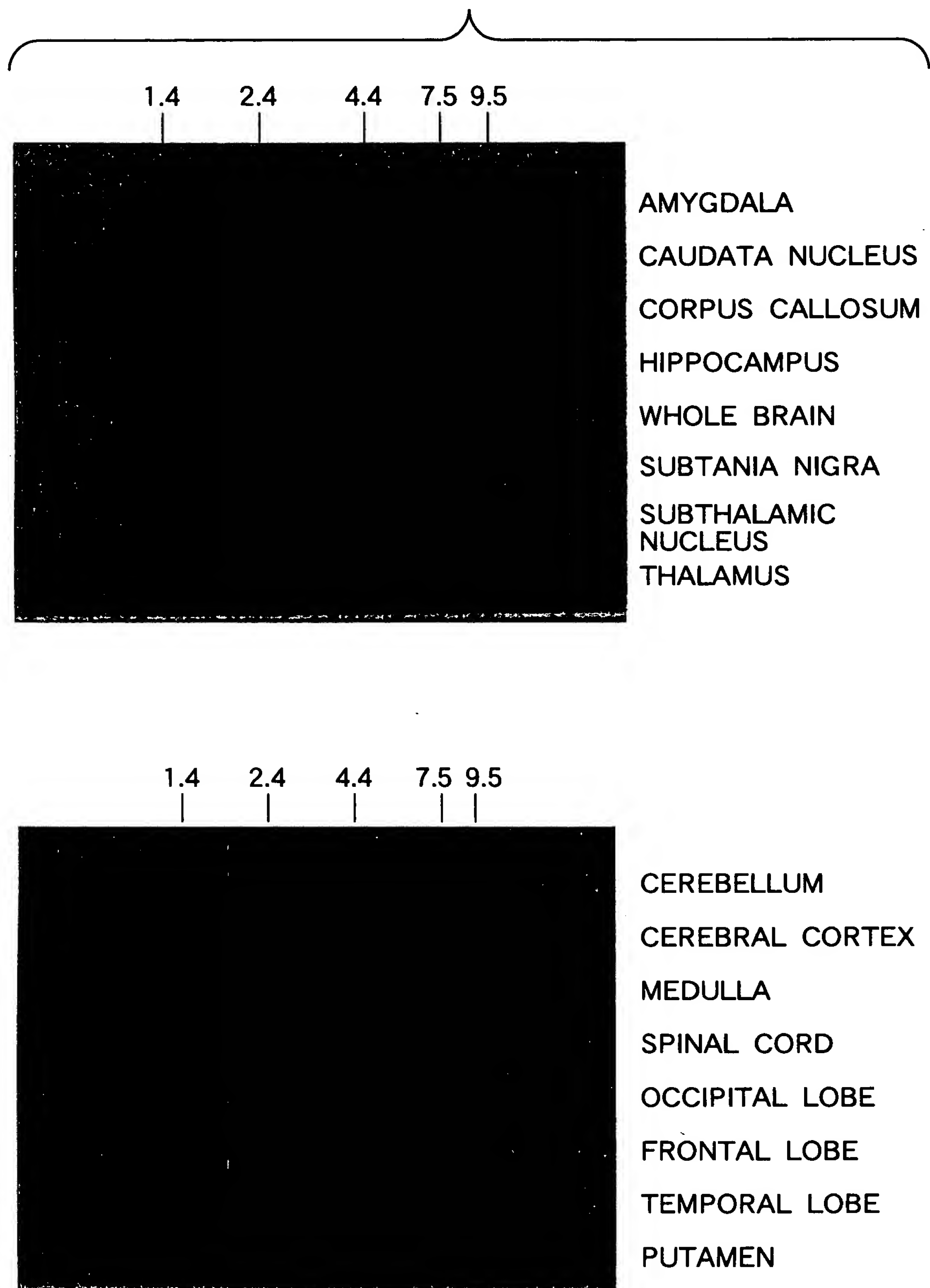


FIG.6

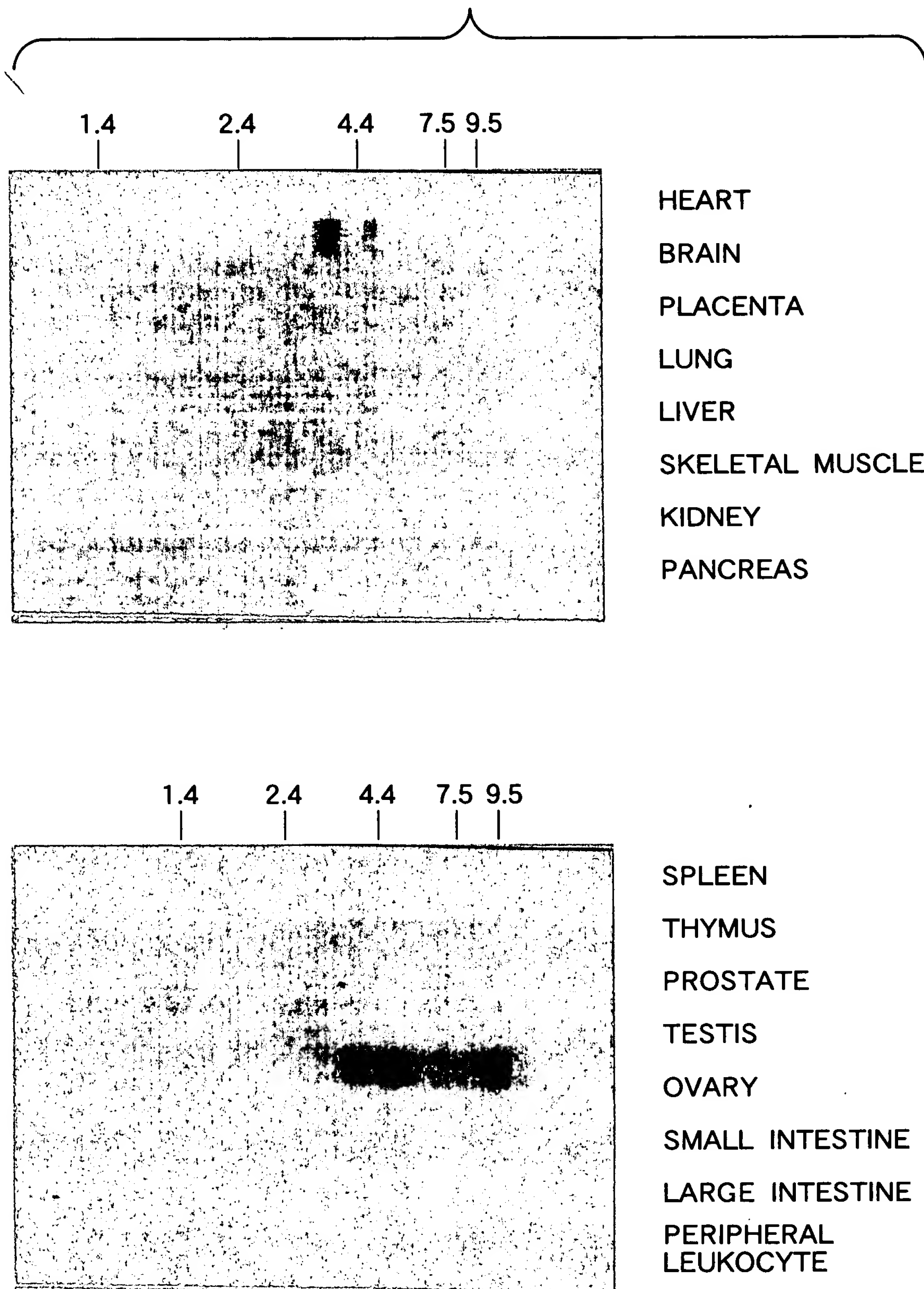


FIG.7

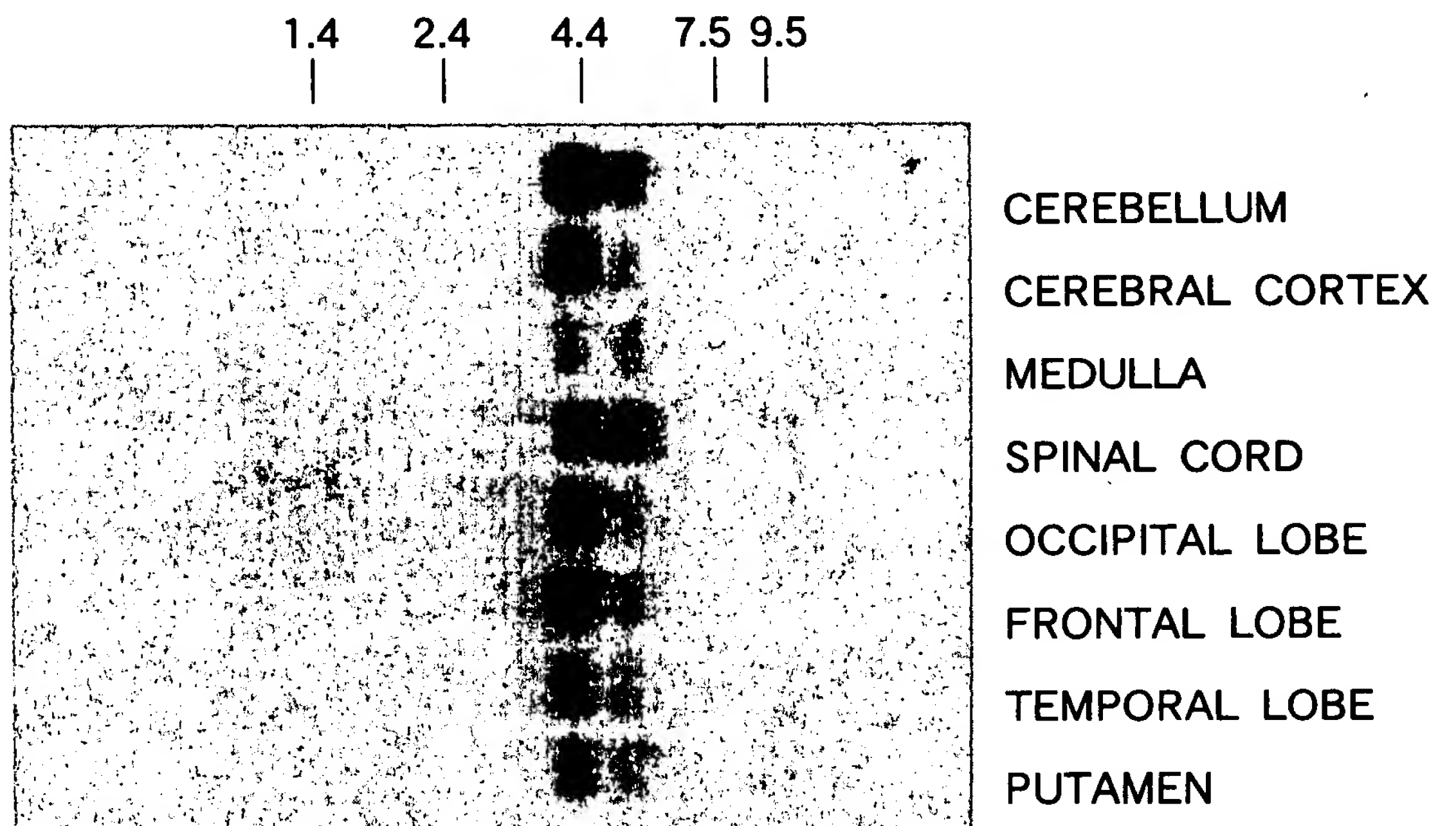
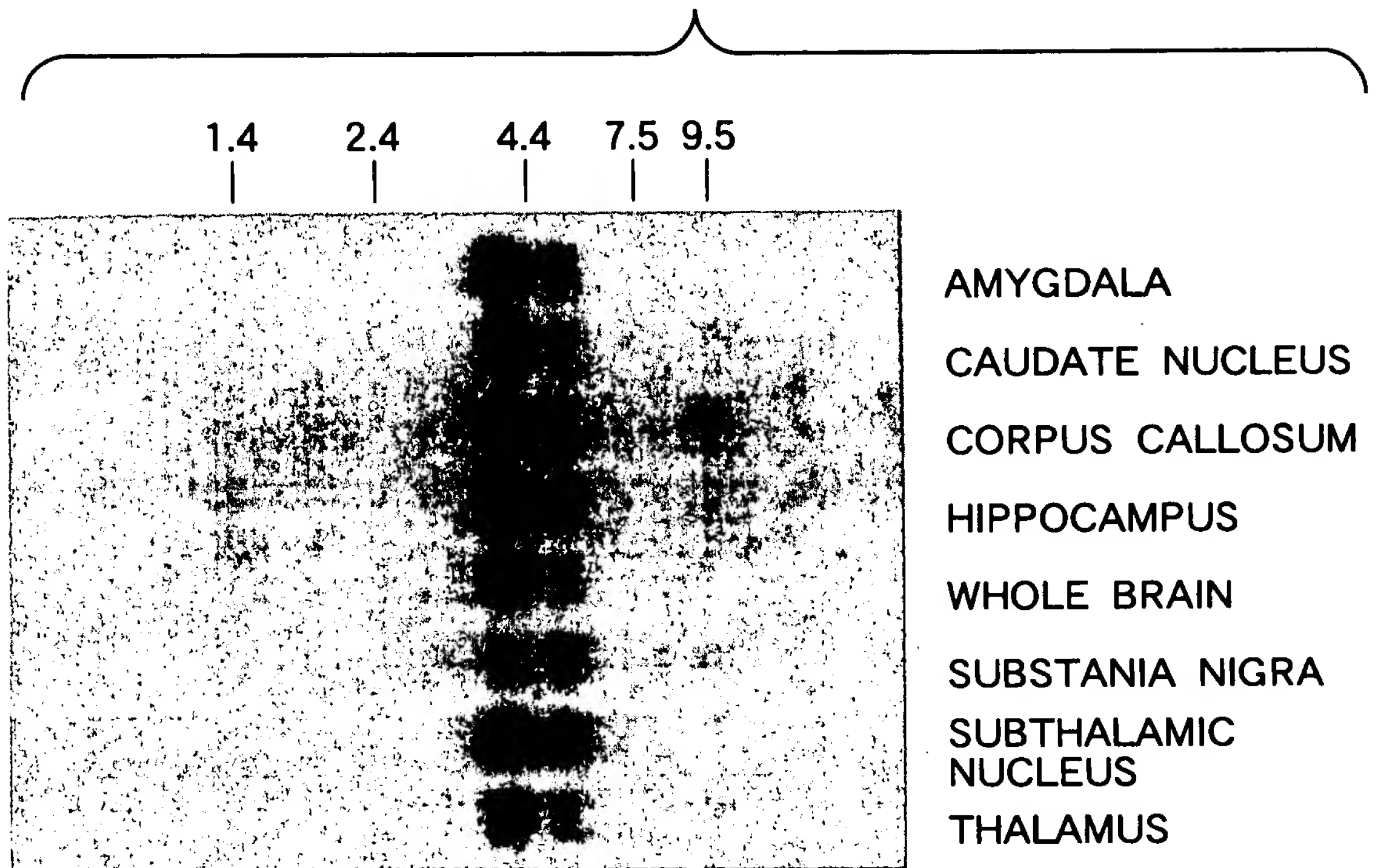


FIG.8

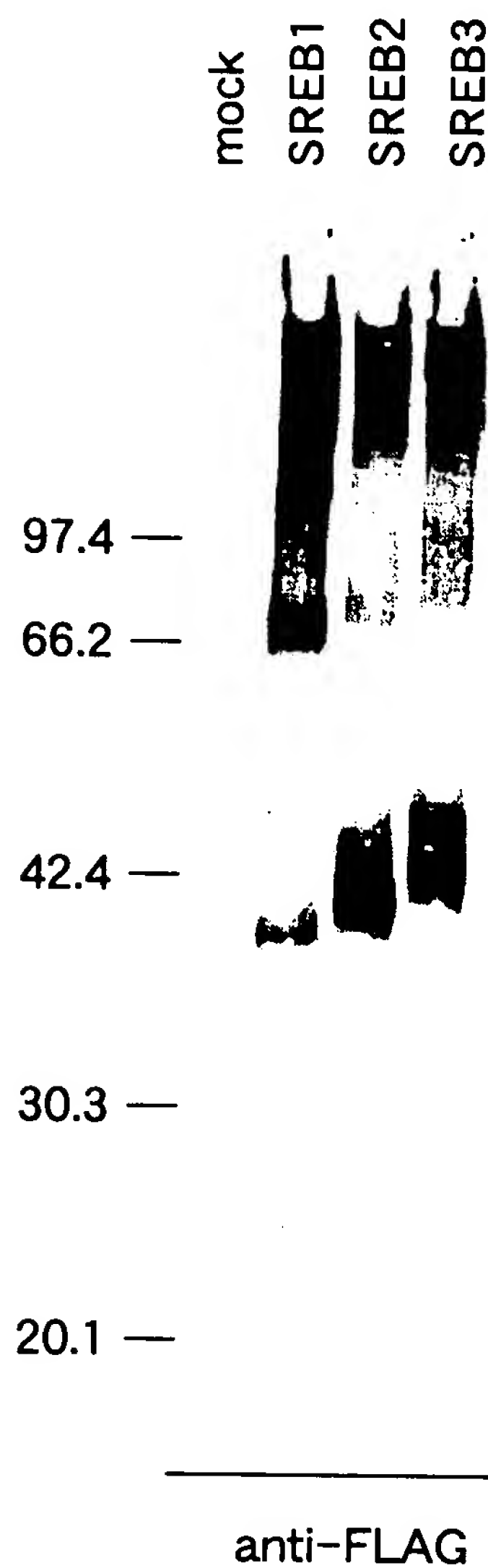


FIG.9

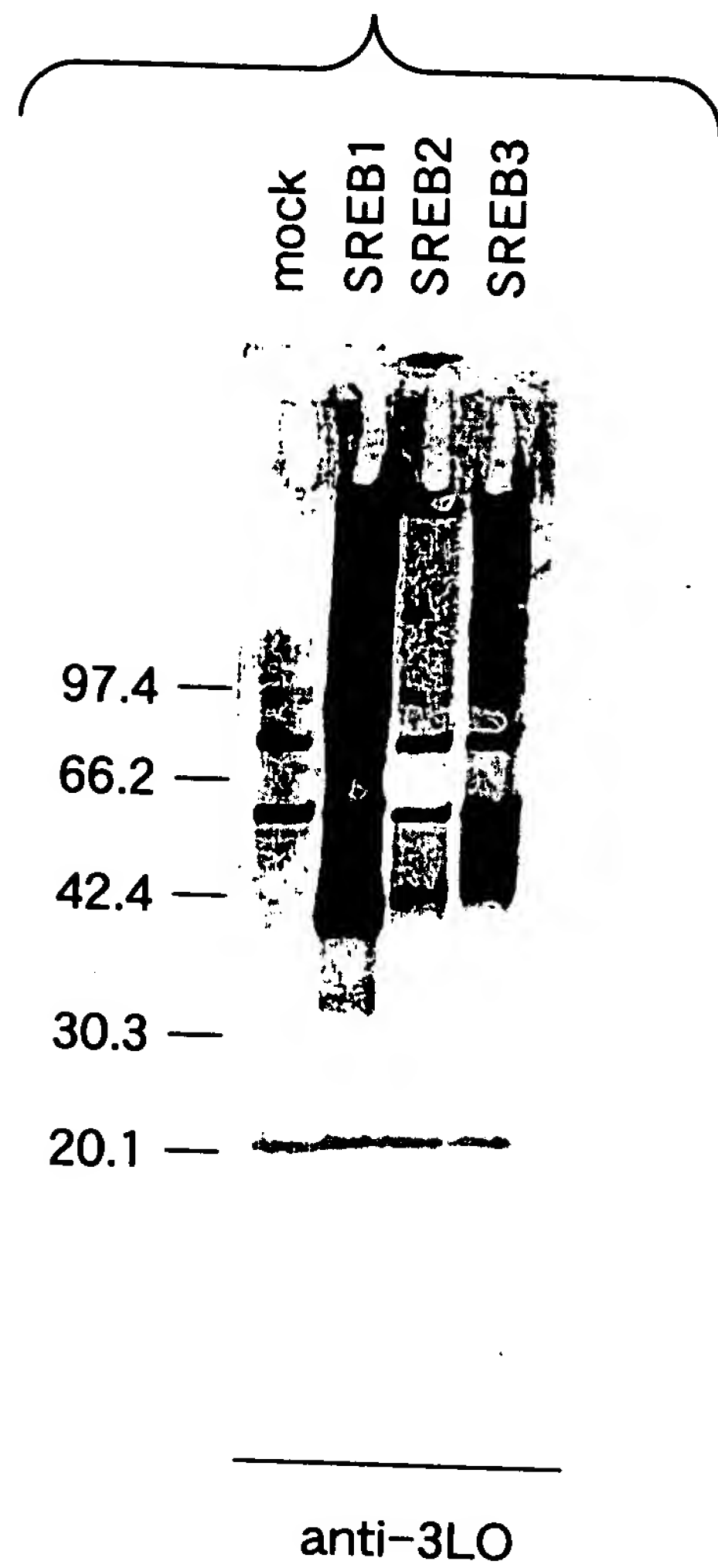


FIG.10

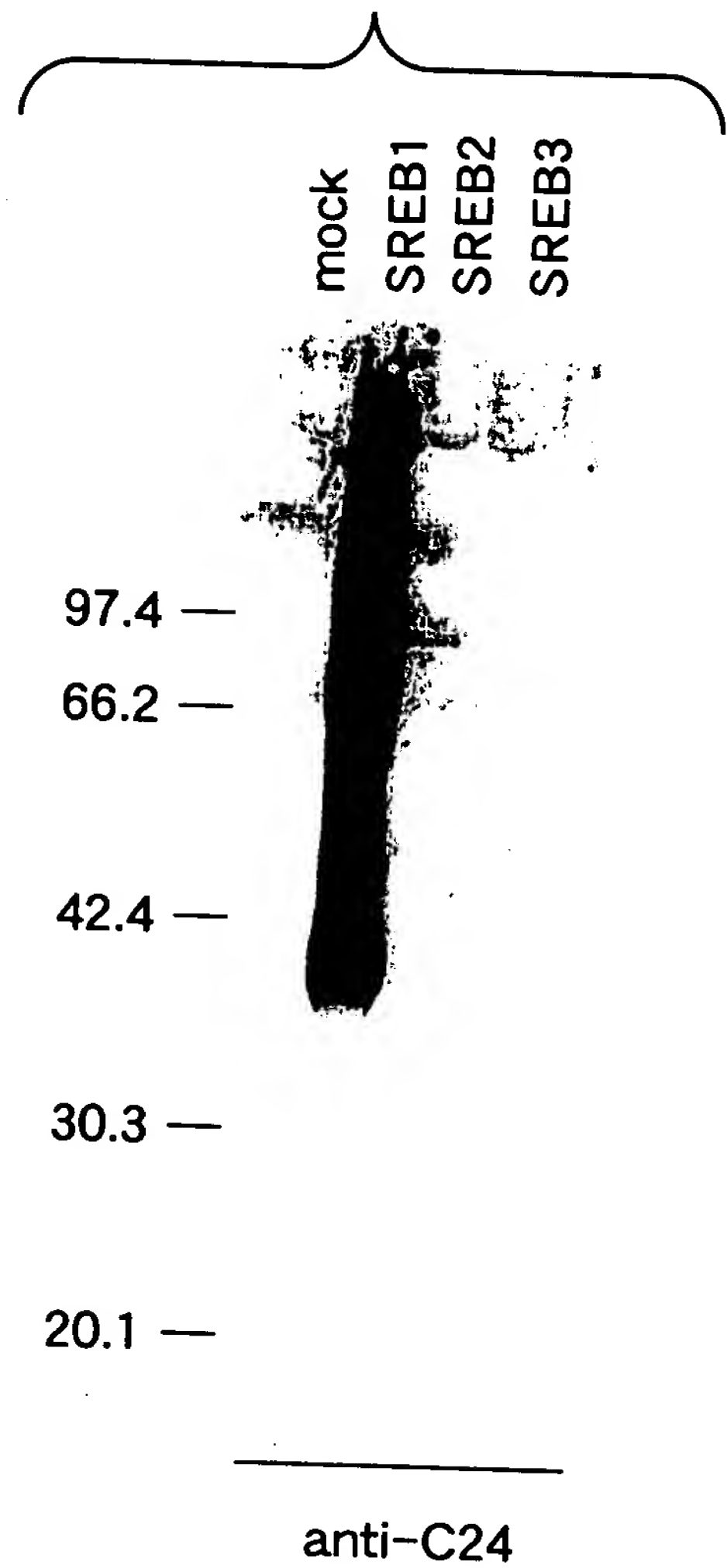


FIG.11

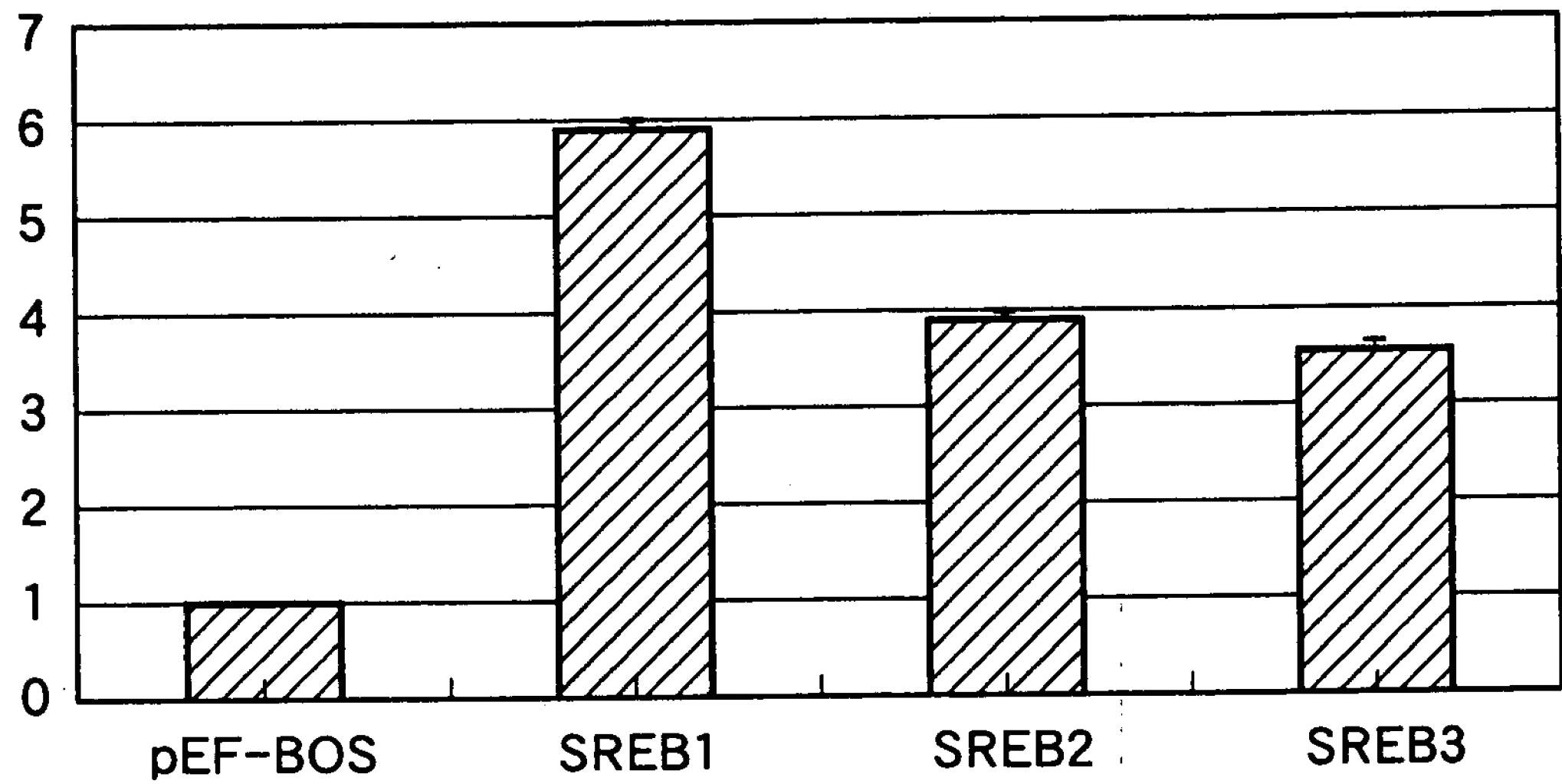


FIG.12

